

Article

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Alketbi, Salem Khalifa and Goodwin, William H

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Salem K. Alketbi, W. Goodwin



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The effect of surface type, collection and extraction methods on Touch DNA

Salem K Alketbi^{1,2*}, Dr W. Goodwin¹

¹University of Central Lancashire, Preston, UK

²General Department of Forensic Science and Criminology, Dubai Police, Dubai, UAE

*Tel: 00447774141205; E-mail: alkitbe.11@hotmail.com

Abstract

There are different variables that affect the success of Touch DNA recovery, including surface type, the collection method used and extraction techniques. This experiment investigated how a range of porous and non-porous surfaces, different DNA collection (cotton swab, nylon flocked swab and SceneSafe Fast™ minitape) and extraction methods (PrepFiler Express BTA™ and QIAamp® DNA Investigator) affected touch DNA recovery.

1. Introduction

Different variables that affect the success of obtaining a substantial quality DNA profile from Touch DNA, including shedder status, surface type, and pressure of contact, as well as the method used for DNA collection and extraction [1]. Although previous studies have investigated the effectiveness of different methods used for DNA recovery, there are some deficiencies of published information regarding touch DNA recovery [2]. The duration, over which the deposit on the surface and the area over which the touch occurs is important, as it helps to evaluate the effectiveness of sampling touch DNA [3]. There is a need to incorporate recovery and extraction efficiencies in the interpretation of trace DNA from various of surfaces [4], and the following study sought to investigate the effect of surface type, DNA collection and extraction methods on Touch DNA recovery.

2. Materials and methods

2.1 Experimental set

A selection of six surfaces (stainless steel, smooth non-porous; glass, smooth non-porous; textured wood, rough porous; banana skin, smooth porous; copier paper, smooth porous and textured plastic, rough non-porous) were chosen to replicate common items encountered in crime scenes. All non-porous surfaces were cleaned with 2% virkon (viricidal disinfectant) and ultraviolet radiation (UV) for 15 min; porous surfaces were irradiated using UV light for 25 min.

2.2 Deposition

Two participants were asked to wash their hands with antibacterial soap and refrain from undertaking any activity for 10 min. Then, they were asked to charge the fingers of both hands with eccrine sweat by touching behind their ears or forehead to load them with DNA. The participants were then asked to touch the surfaces using the index, middle, and ring fingers of both hands separately for deposition by applying medium pressure on a 5 x 7 cm area of the surface for 1 min. The same procedure was repeated on all surfaces for equal deposition. Each participant deposited on separate six surfaces six times to make a total of 72 samples (two replicates for each surfaces), then one participant was asked to do further depositions on glass (n=18) and copier paper (n=18) (three replicates for each surfaces) to test the variables. All the depositions were done at room temperature.

2.3 DNA recovery and extraction

Three different methods were used to recover the touch DNA, Copan cotton swab (150C) (CS), Copan nylon flocked swab (4N6 FLOQSwabs®) (NS), and SceneSafe Fast™ minitape (K545) (MT). Before collection, 100 µL of sterile distilled water was applied to moisten CS using a plastic spray bottle technique (developed in Dubai police forensic DNA lab; each single spray contains approximately 50 µL). For NS, 30 µL of sterile distilled water was applied to moisten the swab using a pipette as recommended by the manufacturer. No water was added to the MT, but to increase the amount of touch DNA collected, each minitape was applied 16 times to the area [5].

Touch DNA was recovered from the surfaces 30 min after deposition and extracted immediately. Samples deposited by the two participants (n=72) were extracted by PrepFiler Express BTA™ kit (Thermo Fisher Scientific) (EXT 1) using an AutoMate Express Forensic DNA Extraction System according to the manufacturer's recommendations and manually using the QIAamp® DNA Investigator Kit (Qiagen) (EXT 2) as per the manufacturers' protocol. However, with EXT 2 nylon swabs were extracted using NAOBasket™ as recommended by Copan to increase the DNA yield.

For the extra deposition made by participants one for glass (n=18) and copier paper (n=18) the same extraction techniques were used, but for EXT 1 460 µL of lysis buffer was used instead of 230 µL. For all extracted samples, full swab heads were used and the lower sticky part of the minitape, with a final extracted sample elution of 50 µL.

2.4 DNA quantification, amplification and analysis

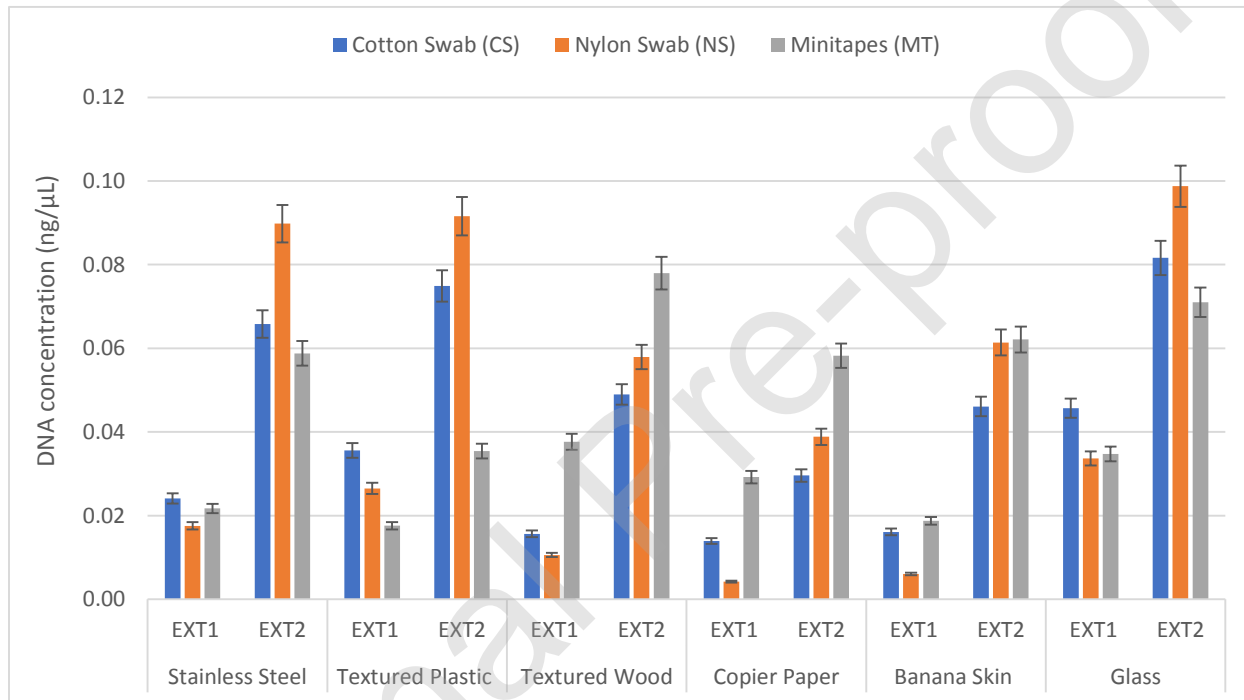
Extracted samples were quantified using the Quantifiler® Human DNA Quantification Kit, QuantStudio 5 Real-Time PCR (qPCR) and HID Real-Time PCR analysis software v1.3 according to the manufacturer's instructions (Thermo Fisher Scientific). Amplification were performed using the GlobalFiler™ PCR amplification Kit (Thermo Fisher Scientific), following the 30 cycles protocol. The data were analysed using GeneMapper® ID-X Software Version 1.2 (Thermo Fisher Scientific). Statistical analysis on the tested variables was performed with RStudio using factorial analysis of variance (ANOVA). In ANOVA the p-value is derived from the F-distribution which is different for every pair of degree of freedom (df) values (F value =variance of the variables means (Mean Square Between) / mean of the within variables variances (Mean Squared Error))

Blanks were taken from surfaces after cleaning, and negative controls for the collection and extraction methods, all of which were DNA-free when quantified. Random samples from the surfaces were amplified to evaluate the quality of samples collected; the amplified samples produced full DNA profiles without any sign of mixtures or contamination.

3. Results and discussion

The amount of DNA collected from the selected six surfaces ($n=72$) was significantly affected by the type of surface ($F_{5, 36} = 3.469$, $p < 0.05$), and the extraction methods ($F_{1, 36} = 72.286$, $p < 0.05$). The highest amount of DNA was recovered from glass surfaces, and the least amount of from copier paper surfaces with both extraction methods. Samples extracted by EXT 2 were higher in DNA than samples extracted by EXT 1 when manufacturer's protocols were followed (Figure 1). That was not the case when 460 μL of lysis buffer were used with EXT 1 instead of 230 μL for the extra samples on glass ($F_{1,12} = 1.05$, $p = 0.33$) and copier paper ($F_{1,12} = 4.74$, $p = 0.05$) (Figure 2).

By analysing the extra samples collected from glass and copier paper individually (Figure 2), the amount of DNA collected from glass ($n=18$) was significantly affected by collection type ($F_{2, 12} = 22.36$, $p < 0.05$), the interaction between collection type and extraction method ($F_{2, 12} = 13.70$, $p < 0.05$). There was not a big difference of DNA extracted by both extraction methods when samples were collected by CS and NS.



In contrary, some collected DNA was not efficiently extracted by EXT 2 when samples were collected by MT (Mean 0.03 ng/μL), when compared to samples collected by MT and extracted by EXT 1 (Mean: 0.06 ng/μL). Furthermore, the amount of DNA collected from copier paper ($n=18$) was significantly affected by collection type ($F_{2,12} = 9.25$, $p < 0.05$) and the interaction between collection type and extraction method ($F_{2,12} = 7.46$, $p < 0.05$). MT was the most suitable collection type for copier paper and EXT 1 the most efficient extraction method with MT (mean: 0.04 ng/μL) compared to EXT 2 (mean: 0.02 ng/μL).

Figure 1. Mean of DNA recovered from the six surfaces ($n = 72$) using three collection methods and extracted using two extraction methods, PrepFiler Express BTA™ kit (EXT 1) and QIAamp® DNA Investigator Kit (EXT 2). CS performed better with EX1 and NS performed better with EXT 2 when used on non-porous surfaces. MT performed better when used on porous surfaces regardless the effect of extraction. Error bars represent 95% confidence intervals.

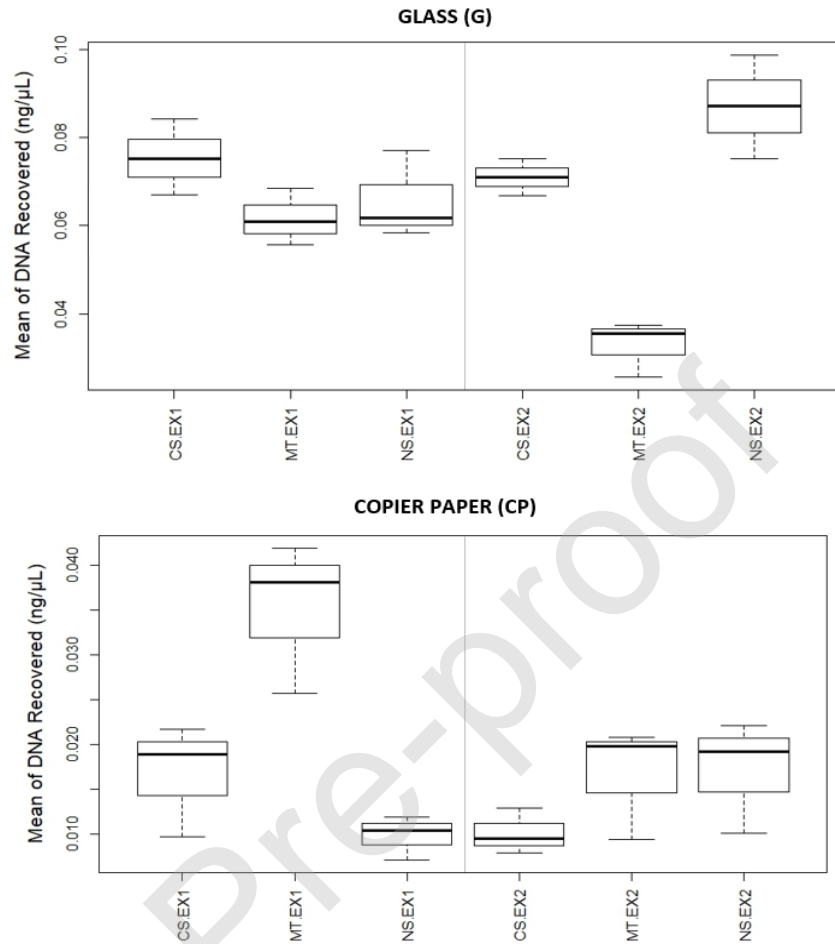


Figure 2 - The mean of DNA recovered from glass (n=18) and copier paper (n=18). Collected by cotton swab (CS), nylon swab (NS) and minitapes (MT), then extracted by two extraction methods [PrepFiler Express BTA™ kit (EX1) and QIAamp® DNA Investigator Kit (EX2)]. Error bars represent standard Error.

4. Conclusion

In conclusion, surfaces retain Touch DNA differently and it is important to consider the most appropriate collection method for different type of surfaces. From the experimental findings, it is recommended to use cotton swabs or nylon swabs for non-porous surfaces such as glass and textured plastic, whereas the use of tapes is better for porous surfaces such as paper or wood. Moreover, for optimal recovery of DNA different types of collection method require different extraction processes. Different extraction methods, automated or manual, have different requirements, with the specimen size in combination with the amount of lysis buffer influencing the maximum DNA yield. Based on these finding, it is recommended to use different protocols to collect and process Touch DNA from various of surfaces.

Conflict of interest

None.

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References

- [1] S.K. Alketbi, The Affecting Factors of Touch DNA. *Journal of Forensic Research*. 9(3) (2018) 424.
- [2] R.J. Brownlow, K.E. Dagnall, C.E. Ames, A comparison of DNA collection and retrieval from two swab types (cotton and nylon flocked swab) when processed using three QIAGEN extraction methods. *Journal of Forensic Sciences*. 57 (3) (2012) 713–717.
- [3] S.H. Tobias, G.S. Jacques, R.M. Morgan, et al., The effect of pressure on DNA deposition by touch. *Forensic Science International: Genetics Supplement Series*. 6 (C) (2017), 12–14.
- [4] D. Taylor, A. Biedermann, L. Samie, et al., Helping to distinguish primary from secondary transfer events for trace DNA. *Forensic Science International: Genetics*. 28 (2017) 155–177.
- [5] T.J. Verdon, R.J. Mitchell, R.A. van Oorschot, Evaluation of tapelifting as a collection method for touch DNA. *Forensic Science International: Genetics*. 8(1) (2014) 179–186.